

**Fig. 1**





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**PANTHER Prowler** / Microsoft Internet Explorer

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Address <http://panther.appliedbiosystems.com/>

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**PANTHER**  
Classification System

PANTHER gene list Customize gene list

checked items Display Gene List Send To Workspace

Display: 10 UPDATE Expanded View Filtered By: Panther Molecular Function FILTER Advanced Filter

Hits 1-10 of 5414 [page: (1) 2 3 4 5 6 7 8 9 10 >>]

Gene ID	Gene Name	Hit family (CF#) or subfamily (SF#)	Panther best Score/Relation	Panther Molecular Function	Panther Biological Processes	GeneXAssay
1. LocusID:13424	dynein, cytoplasmic, heavy chain1 Dnchc1	View unavailable to non-CDS users	View unavailable to non-CDS users	Microtubule binding motor protein	General vesicle transport	Mm00466548 m
2. LocusID:29489	dynein, cytoplasmic, heavy chain 1 Dnchc1	View unavailable to non-CDS users	View unavailable to non-CDS users	Microtubule binding motor protein	General vesicle transport	Rn00570138 m1
3. CG7507	dynein heavy chain 64C Dhc64c	View unavailable to non-CDS users	-6740.02 ●●●	Microtubule binding motor protein	General vesicle transport	
4. LocusID:65209	dynein, cytoplasmic, heavy chain 2 Dnch2	View unavailable to non-CDS users	View unavailable to non-CDS users	Microtubule binding motor protein	General vesicle transport	Rn00576479 m1
5. LocusID:15194	Huntington disease gene	View unavailable to non-CDS users	View unavailable to non-CDS	Molecular function unclassified	General vesicle transport Other neuronal activity	Hs00169273 m1

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Fig. 3

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PANTHER Proviewer - Microsoft Internet Explorer

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Advanced Filter

0 UPDATE

GeneXAssay	SNP Assay	Celera SNP ID	Transcript ID	Protein ID	Celera Start Pos	Celera End Pos	Celera Location (chromosome)	Public Start Pos	Public End Pos	Public Location (chromosome)	Species
Mm00466548 m1			XM 109305 (2 transcripts)	XM 109305 (2 proteins)	View unavailable to non-CDS users	View unavailable to non-CDS users	View unavailable to non-CDS users	132372166	132425699	12	NCBI: M. musculus
Rn00570138 m1			NM 019226	NP 062099	View unavailable to non-CDS users	View unavailable to non-CDS users	View unavailable to non-CDS users	4161977	4400166	8	NCBI: R. norvegicus
Rn00576479 m1			CG7507-RA (2 transcripts)	CG7507-PA (2 proteins)	View unavailable to non-CDS users	View unavailable to non-CDS users	View unavailable to non-CDS users	33033473	33181273	5	NCBI: M. musculus

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Fig. 4



**Fig. 5**

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**PANTHER**  
Classification System

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PANTHER transcript list Customize transcript list

Checked items: Display Transcript/Protein Send To Workspace

Display: 10 UPDATE Filtered By: Protein ID FILTER HMM Score Cutoff (<): 0.0 UPDATE

Hits 1-10 of 5414 [page: (1) 2 3 4 5 6 7 8 9 10 >>]

Transcript ID	Protein ID	Gene ID	Gene Name	Hit family (CF#) or subfamily (SF#)	Panther best Score/Relation	Panther Molecular Function	Panther Biological Processes	GeneXAssav
<input type="checkbox"/> 1. XM 109305	XP 109305	LocusID:13424	dynein, cytoplasmic, heavy chain 1 Dnchc1	View unavailable to non-CDS users	View unavailable to non-CDS users	Microtubule binding motor protein	General vesicle transport	
<input type="checkbox"/> 2. NM 030238	NP 084524	LocusID:13424	dynein, cytoplasmic, heavy chain 1 Dnchc1	View unavailable to non-CDS users	View unavailable to non-CDS users	Microtubule binding motor protein	General vesicle transport	Mm00466548
<input type="checkbox"/> 3. NM 019226	NP 062099	LocusID:29489	dynein, cytoplasmic, heavy chain 1 Dnchc1	View unavailable to non-CDS users	View unavailable to non-CDS users	Microtubule binding motor protein	General vesicle transport	Rn00570138
<input type="checkbox"/> 4. CG7507-RA	CG7507-PA	CG7507	Dynein heavy chain 64C Dnc64C	CYTOPLASMIC DYNEIN HEAVY CHAIN [CF12238-SF9]	-6740.02 •••	Microtubule binding motor protein	General vesicle transport	
<input type="checkbox"/> 5. NM 023024	NP 075413	LocusID:65209	dynein, cytoplasmic,	View unavailable to non-CDS	View unavailable to non-CDS	Microtubule binding motor protein	General vesicle transport	Rn00576479

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Fig. 6

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List Send To Workspace

Filtered By: Protein ID FILTER Advanced Filter

HMM Score Cutoff (<): 0.0 UPDATE

Name	Hit	Panther Score/Relation	Panther Molecular Function	Panther Biological Processes	SNP Assay	Celera SNP ID	Species
In, asmic, chain1	View unavailable to non-CDS users	View unavailable to non-CDS users	Microtubule binding motor protein	General vesicle transport			NCBI: M. Musculus
In, asmic, chain 1	View unavailable to non-CDS users	View unavailable to non-CDS users	Microtubule binding motor protein	General vesicle transport	Mm00466548 m1		NCBI: M. musculus
In, asmic, chain 1	View unavailable to non-CDS users	View unavailable to non-CDS users	Microtubule binding motor protein	General vesicle transport	Rn00570138 m1		NCBI: R. norvegicus
In heavy 84C 4C	View unavailable to non-CDS users	View unavailable to non-CDS users	Microtubule binding motor protein	General vesicle transport			FlyBase: D. melanogaster
In, asmic, chain1	View unavailable to non-CDS users	View unavailable to non-CDS users	Microtubule binding motor protein	General vesicle transport	Rn00576479 m1		NCBI: R. norvegicus

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Fig. 7

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Back Address http://bobcat.fc.celera.com:6677/prowler/index\_new.jsp Go

**Celera Discovery System**

BioMolecule Library Text/Sequence Analysis Workspace Help Logout

**Panther Protein Family Viewer** SEARCH: Categories families/subfamilies **apoptosis** go

SPECIES: ☒ *H. sapiens* ☒ *M. musculus* ☒ *D. melanogaster* Use Case Tips

Categories: ☐ or ☐ and update families clear Families: update categories gene list clear

- SIGNAL TRANSDUCTION
- INTRACELLULAR PROTEIN TRAFFIC
- PROTEIN TARGETING AND LOCALIZATION
- TRANSPORT
- IMMUNITY AND DEFENSE
- ONCOGENESIS
- NEURONAL ACTIVITIES
- MUSCLE CONTRACTION
- BLOOD CLOTTING
- HOMEOSTASIS
- SENSORY PERCEPTION
- DEVELOPMENTAL PROCESSES
- CELL CYCLE
- BLOOD CIRCULATION AND GAS EXCHANGE
- APOPTOSIS**
- CELL STRUCTURE AND MOTILITY

- CALPAIN THIOLESTERASE C2 CF10557
  - Family tree Full MSA Partial MSA
- RELAXIN CF12004
  - Family tree Full MSA Partial MSA
- WD DOMAIN-CONTAINING PROTEIN CF11554
  - Family tree Full MSA Partial MSA
- CASPASE-RELATED CF10454
  - Family tree Full MSA Partial MSA
- INTERLEUKIN 6 CF11457
  - Family tree Full MSA Partial MSA
- TRANSCRIPTION FACTOR ETS-RELATED CF11849
  - ETS-RELATED PROTEIN (SF11)
  - ETS DNA-BINDING PROTEIN (SF2)
  - ETS-RELATED PROTEIN TEL (SF6)
  - TRANSCRIPTION FACTOR TEL 2 (SF4)
  - C-ETS-2-RELATED (SF1)
  - gb def: lin-1 (caenorhabditis elegans) (SF0)
  - ETS DOMAIN PROTEIN ELK-4 (SF13)

**Fig. 8**

TABLE 1 • Biological function enrichments in cell-cycle-regulated expression clusters

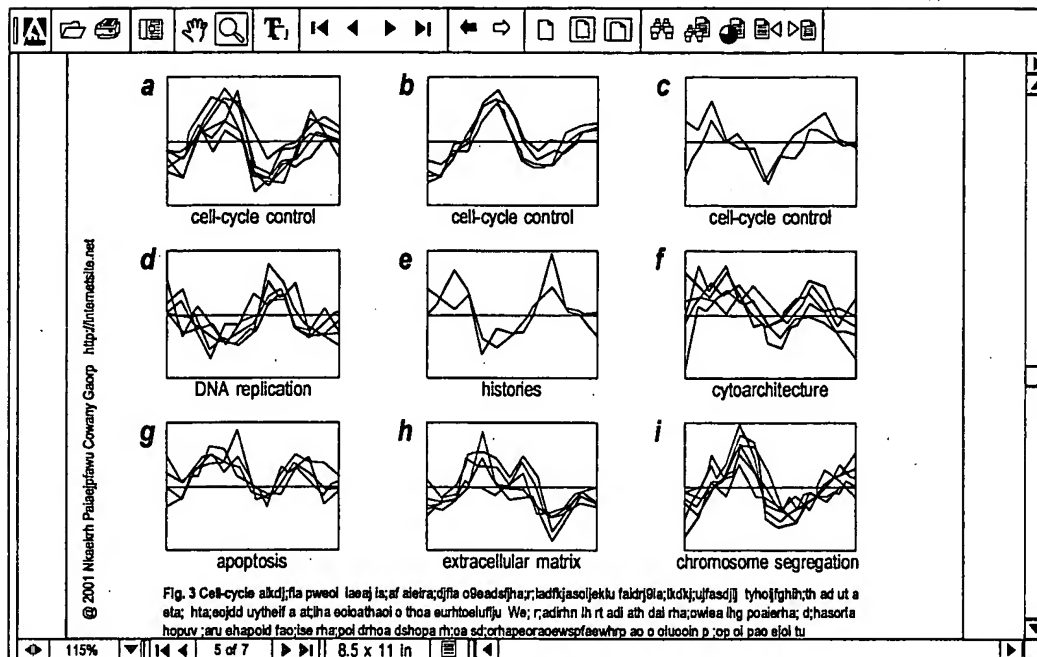
Biological function	late G <sub>1</sub> (53)	S (107)	G <sub>2</sub> (108)	M (119)
amino acid metabolism (35)	0.1 0/0	0.5 1/1	0.5 1/1	0.3 0/1
cell-to-cell adhesion (137)	0.5 0/1	0.7 3/2	0.6 1/2	4.7 11/2
chromosome segregation (17)	0.1 0/0	0.1 0/0	2.7 3/0	5.0 5/0
cytokine signaling (151)	0.5 0/1	0.9 4/2	0.6 2/2	1.1 0/3
cytoskeletal reorganization (33)	0.1 0/0	1.9 3/0	6.1 7/1	0.5 1/1
DNA replication (47)	7.2 7/0	0.3 0/1	0.3 0/1	0.8 2/1
glycolysis (31)	0.1 0/0	0.2 0/0	0.2 0/0	0.2 0/1
G-protein signaling (223)	0.7 0/2	0/7 3/3	1.0 1/3	0.7 4/4
immune regulation (274)	0.6 2/2	1.2 1/4	1.2 1/4	1.0 2/5
intracellular transport (114)	0.4 1/1	1.2 4/2	0.5 1/2	0.8 3/2
ionic homeostasis (47)	0.6 1/0	0.3 0/1	0.3 0/3	0.3 0/1
mitosis and cell-cycle control (89)	0.3 12/1	0.5 1/1	5.0 8/1	5.7 10/1
mRNA regulation (553)	1.3 7/4	0.9 10/8	1.0 8/8	0.9 8/9
muscular contraction (82)	0.3 0/0	2.1 5/1	5.3 9/7	0.3 1/1
neurotransmitter signaling (69)	0.2 0/1	0.5 0/1	0.4 1/1	0.5 0/1
PIP signaling (51)	0.2 0/0	0.3 0/1	0.4 1/1	0.4 1/1
protein phosphorylation (292)	0.6 2/2	0.7 4/4	1.1 7/4	1.0 7/5
translation (102)	1.5 3/1	0.5 1/2	0.7 0/2	0.8 0/2

Shown are representative biological a;dkh ad; a;olehatowiera a;shoahfoacowpoijola ueorha faeuew radscola ueworhpo a;tksdjipa anoeija oiejapo

**Fig. 9**



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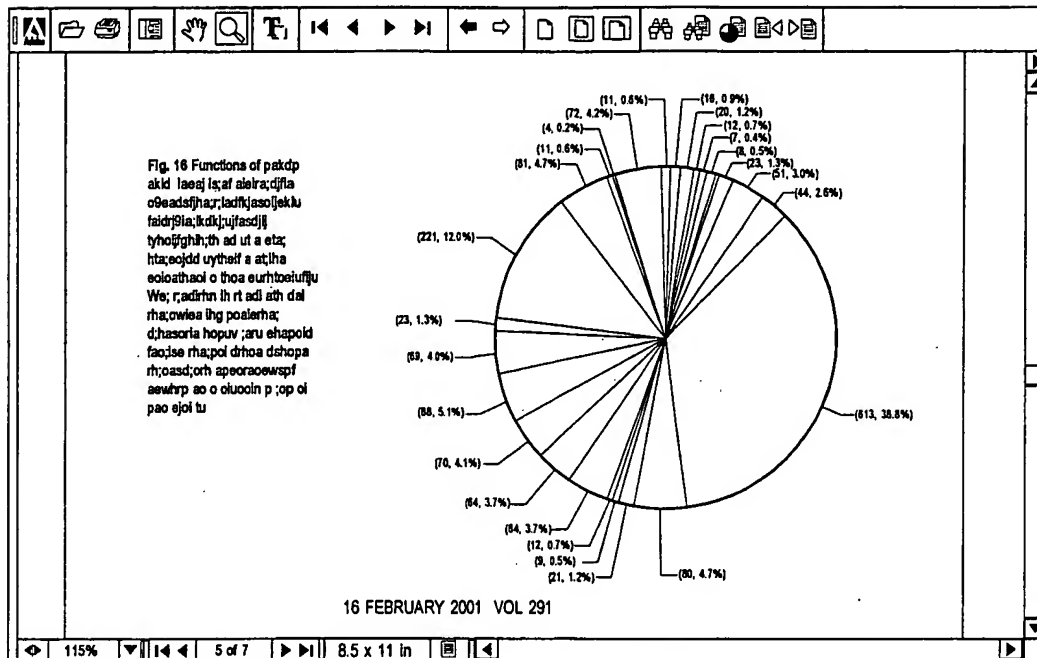
**Fig. 10**

Figure 11 displays a web browser window showing a table of protein families and their counts across four species: H. sapiens (H), D. melanogaster (F), C. elegans (W), S. cerevisiae (Y), and A. thaliana (A). The table is titled "TABLE 19. Number of proteins assigned to selected Panther families or subfamilies in H. sapiens (H), D. melanogaster (F), C. elegans (W), S. cerevisiae (Y), and A. thaliana (A).". The browser interface includes a toolbar at the top and a status bar at the bottom showing "115%" zoom and "5 of 7" pages.

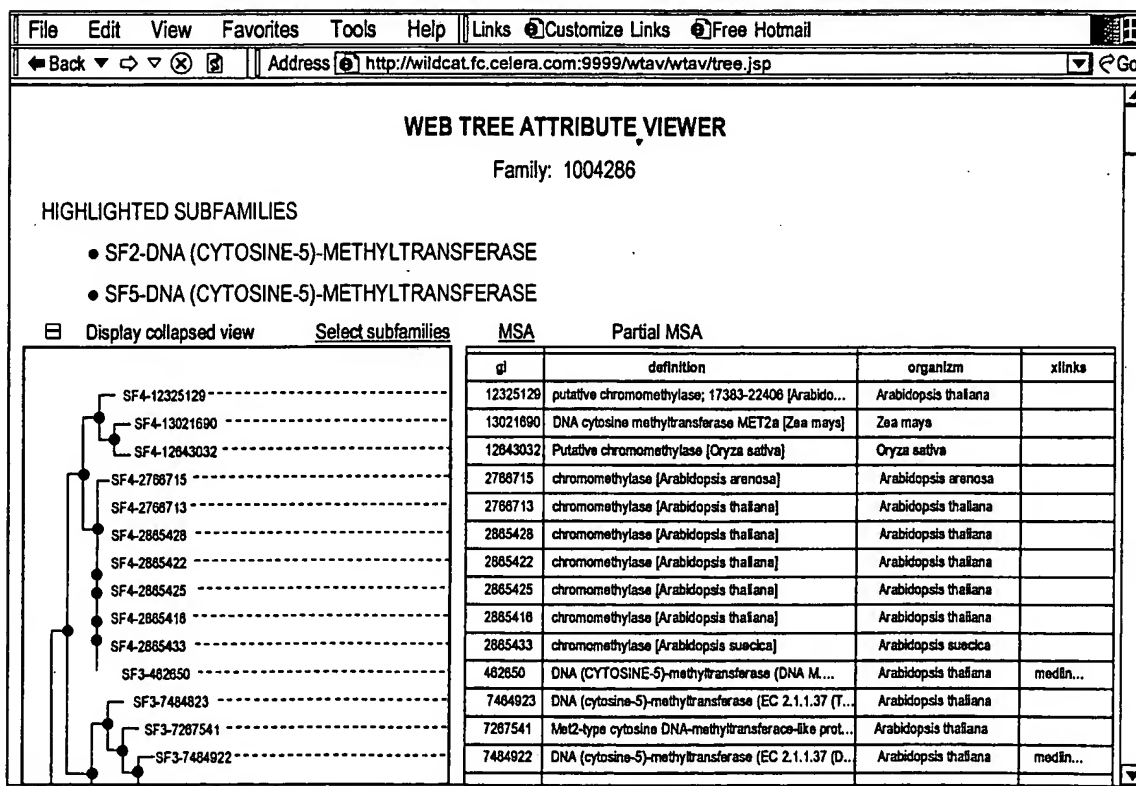
Panther family/subfamily	H	F	W	Y	A
Neural structure, function, development					
Ependymin	1	0	0	0	0
Ion Channels					
Acetylcholine receptor	17	12	56	0	0
Amiloride-sensitive/degenerin	11	24	27	0	0
CNG/EAG	22	9	9	0	30
IRK	16	3	3	0	0
ITP/ryanodine	10	2	4	0	0
Neurotransmitter-gated	61	51	59	0	19
P2X purinoceptor	10	0	0	0	0
TASK	12	12	48	1	5
Transient receptor	15	3	3	1	0
Voltage-gated Ca <sup>2+</sup> alpha	22	4	8	2	2
Voltage-gated Ca <sup>2+</sup> alpha-2	10	3	2	0	0
Voltage-gated Ca <sup>2+</sup> beta	5	2	2	0	0
Voltage-gated Ca <sup>2+</sup> gamma	1	0	0	0	0
Voltage-gated K <sup>+</sup> alpha	33	5	11	0	0
Voltage-gated KQT	6	2	3	0	0
Voltage-gated Na <sup>+</sup>	11	4	4	9	1
Myelin basic protein	1	0	0	0	0
Myelin PO	5	0	0	0	0
Myelin proteolipid	3	1	0	0	0
Myelin-oligodendrocyte glycoprotein	1	0	0	0	0
Neuropilin	2	0	0	0	0
Plexin	9	2	0	0	0
Semaphorin	22	6	2	0	0
Synaptotagmin	10	3	3	0	0

**Fig. 11**

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**Fig. 12**



**Fig. 13**

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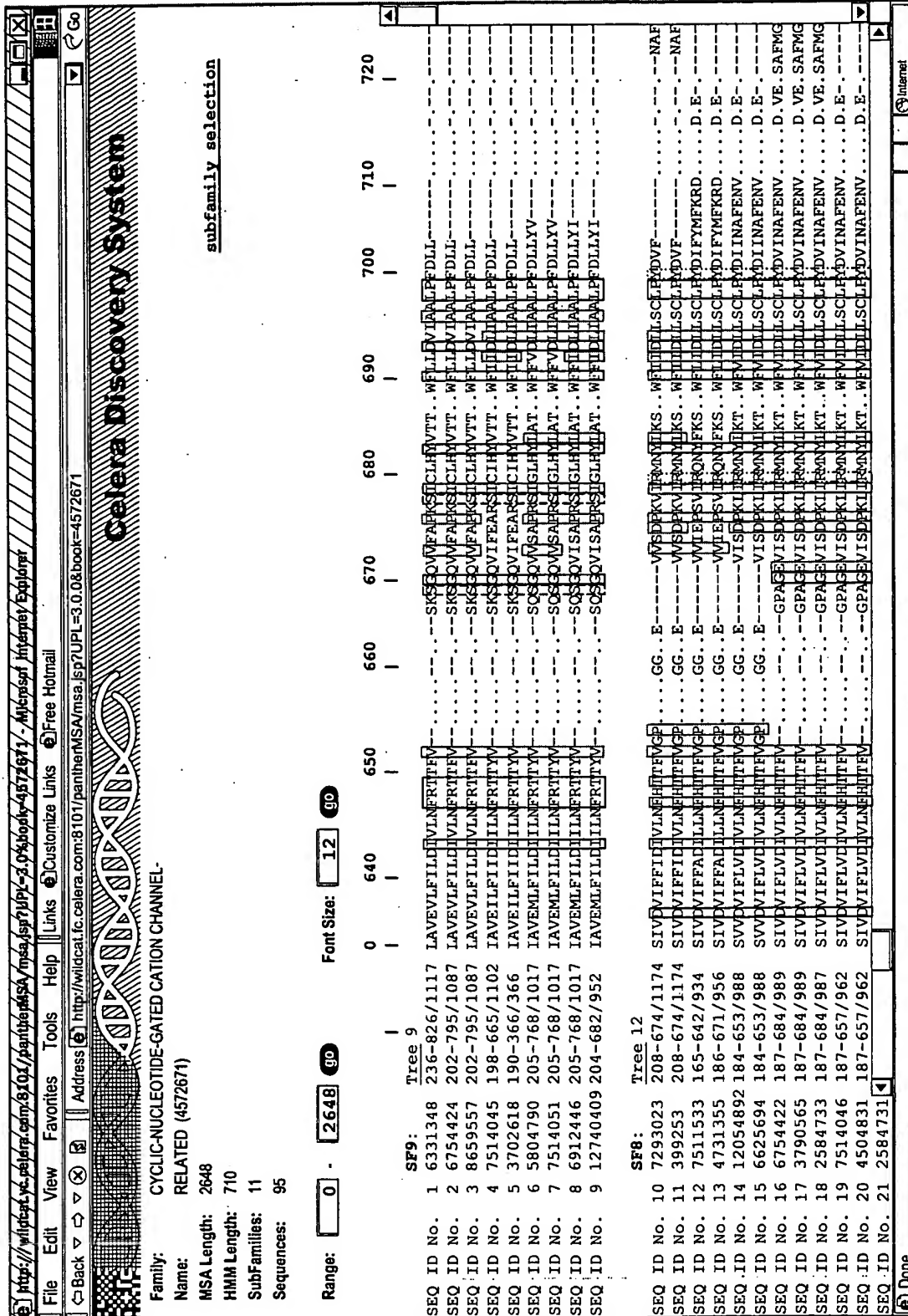



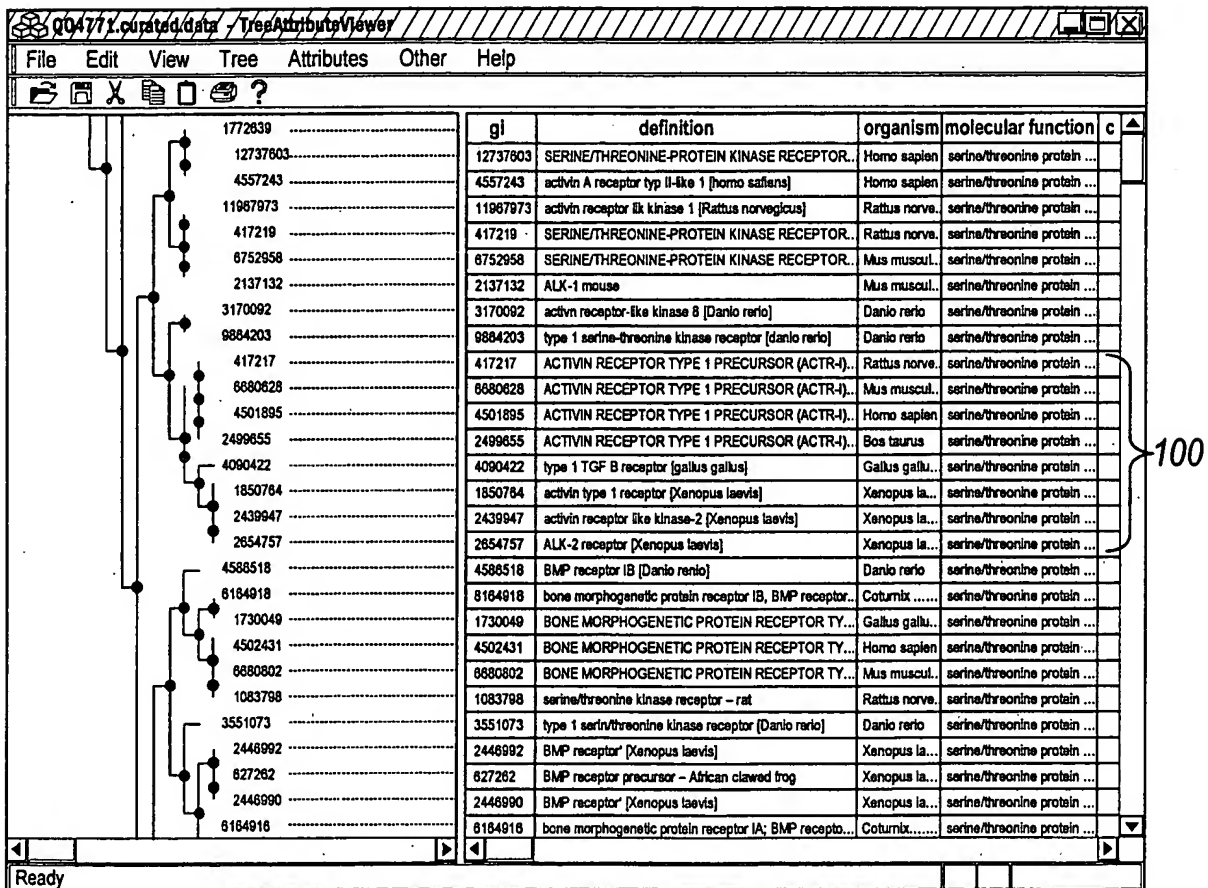
Fig. 14

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PANTHER Discovery Zone		
Panther Classification		
Protein Sequence: <u>CP39928</u>		
Family	Subfamily	SAM NLL- NULL score
<u>CYCLIC-NUCLEOTIDE-GATED CATION CHANNEL-RELATED (2129627)</u> 	<u>gb def:(ae003455) cql7922 gene product [drosophila melanogaster] (2129627SF1)</u>	-875.46
	<u>CYCLIC-NUCLEOTIDE-GATED CATION CHANNEL (2129627:SF2)</u>	-390.97
	<u>CYCLIC-NUCLEOTIDE-GATED CATION CHANNEL (2129627:SF7)</u>	-340.45
	<u>CYCLIC-NUCLEOTIDE-GATED CATION CHANNEL (2129627:SF9)</u>	-301.98
	Family Level Hit	-235.26


**Fig. 15**

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**Fig. 16**

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**Panther Protein  
Function - Family Browser**  
[Tips](#)

SEARCH: ☒ Categories ☒ Families

SPECIES (gene list): ☒ H. sapiens ☒ M. musculus ☒ D. melanogaster

Categories:  ☒ or ☒ and

Families:

1. Select Categories ☐ below.

2. Click on "Update Families" button to view associated data.

Selecting the '+' expands and '-' collapses categories.


- Molecular Functions - 0
- Biological Processes - 15
  - Carbohydrate metabolism
  - Amino-acid metabolism
  - Lipid, fatty acid and steroid metabolism
    - Fatty acid metabolism
    - Steroid metabolism
    - Lipid metabolism
    - Phospholipid metabolism
    - Lipid and fatty acid binding
    - Regulation of lipid, fatty acid and steroid meta
    - Lipid and fatty acid transport
    - Other lipid, fatty acid and steroid metabolism
  - Nucleoside, nucleotide and nucleic acid metabolism
  - Protein metabolism and modification

1. Select families ☐ and or subfamilies ☐ below.

2. Click on "Go to Genelist" button to view associated genes.

Highlighted subfamilies correspond to matches with your selected categories.

**Fig. 17**

**Panther Protein  
Function - Family Browser**  
[Tips](#)

SEARCH: ☒ Categories ☒ Families

SPECIES (gene list): ☒ H. sapiens ☒ M. musculus ☒ D. melanogaster

Categories:  ☒ or ☒ and

Families:

1. Select Categories ☐ below.

2. Click on "Update Families" button to view associated data.

Selecting the '+' expands and '-' collapses categories.

- Molecular Functions - 0
- Biological Processes - 15
  - Carbohydrate metabolism
  - Amino-acid metabolism
  - Lipid, fatty acid and steroid metabolism
    - Fatty acid metabolism
    - Steroid metabolism
    - Lipid metabolism
    - Phospholipid metabolism
    - Lipid and fatty acid binding
    - Regulation of lipid, fatty acid and steroid meta
    - Lipid and fatty acid transport
    - Other lipid, fatty acid and steroid metabolism
  - Nucleoside, nucleotide and nucleic acid metabolism
  - Protein metabolism and modification

1. Select Families ☐ and or subfamilies ☐ below.



2. Click on "Go to Genelist" button to view associated data.

Highlighted subfamilies correspond to searches with your selected categories

- APOLIPOPROTEIN - CF11428 - [7/8]  
Family tree Full MSA Partial MSA
- DIPHOSPHOMEVALONATE DECARBOXYLASE-RELATED - CF10977 [1/20]  
Family tree Full MSA Partial MSA
- FATTY ACID DESATURASE - CF10486 [5/5]  
Family tree Full MSA Partial MSA
- N-ACETYLGLUCOSAMINYL TRANSFERASE COMPONENT GPI1-RELATED  
Family tree Full MSA Partial MSA
- ACYL-COENZYME A OXIDASE-RELATED - CF11520 [15/20]  
Family tree Full MSA Partial MSA
- PATCHED PROTEIN-RELATED - CF10482 [2/8]  
Family tree Full MSA Partial MSA
- OXYSTEROL BINDING PROTEIN-RELATED - CF10972 [7/10]  
Family tree Full MSA Partial MSA
- 3 BETA-HYDROXYSTEROID DEHYDROGENASE-RELATED - CF11580 [8/1]

**Fig. 18**

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<div>  <b>Function Family Browser:</b> Panther Gene List  <div>HMM Score Cutoff (&lt;): <input type="text" value="-35.0"/> Display: <input type="text" value="30"/> <input type="button" value="update"/> <input type="button" value="export"/></div> </div>						
• Sort results by selecting column title. Columns will sort in descending or ascending order.						
<input type="button" value="clr"/> <input checked="" type="button" value="all"/>	ID-protein	Panther Best Hit – Panther ID family (CF#) or subfamily (SF#)	Panther Score/Relation			Species
<input type="checkbox"/> 345.	gene1	ORPHAN NUCLEAR HORMONE RECEPTOR LRH (CF11154:SF208)	-641.6	...		<i>H. sapiens</i>
<input type="checkbox"/> 346.	gene2	ORPHAN NUCLEAR HORMONE RECEPTOR LRH (CF11154:SF208)	-641.51	...		<i>H. sapiens</i>
<input type="checkbox"/> 347.	gene3	RETINOIC ACID RECEPTOR RXR-ALPHA (CF11154:SF218)	-670.15	...		<i>H. sapiens</i>
<input type="checkbox"/> 348.	gene4	RETINOIC ACID RECEPTOR RXR-GAMMA (CF11154:SF217)	-623.88	...		<i>H. sapiens</i>
<input type="checkbox"/> 349.	gene5	RETINOIC ACID RECEPTOR RXR-BETA (CF11154:SF218)	-611.24	...		<i>H. sapiens</i>
<input type="checkbox"/> 350.	gene6	PHOSPHOLIPASE D1 (CF11198:SF4)	-2100.67	...		<i>H. sapiens</i>
<input type="checkbox"/> 351.	gene7	PHOSPHOLIPASE D2 (CF11198:SF5)	-2047.31	...		<i>H. sapiens</i>
<input type="checkbox"/> 352.	gene8	INOSITOL PHOSPHATASE SKIP (CF11200:SF21)	-361.22	...		<i>H. sapiens</i>
<input type="checkbox"/> 353.	gene9	INOSITOL PHOSPHATASE SKIP (CF11200:SF21)	-343.71	...		<i>H. sapiens</i>
<input type="checkbox"/> 354.	gene10	INOSITOL PHOSPHATASE SKIP-RELATED (CF11200:SF22)	-389.15	...		<i>D. melanogaster</i>
<input type="checkbox"/> 355.	gene11	INOSITOL PHOSPHATASE SKIP-RELATED (CF11200:SF22)	-354.86	...		<i>D. melanogaster</i>
<input type="checkbox"/> 356.	gene12	SPHINGOSINE PHOSPHATE LYASE-RELATED (CF11253:SF10)	-722.67	...		<i>D. melanogaster</i>

**Fig. 19**





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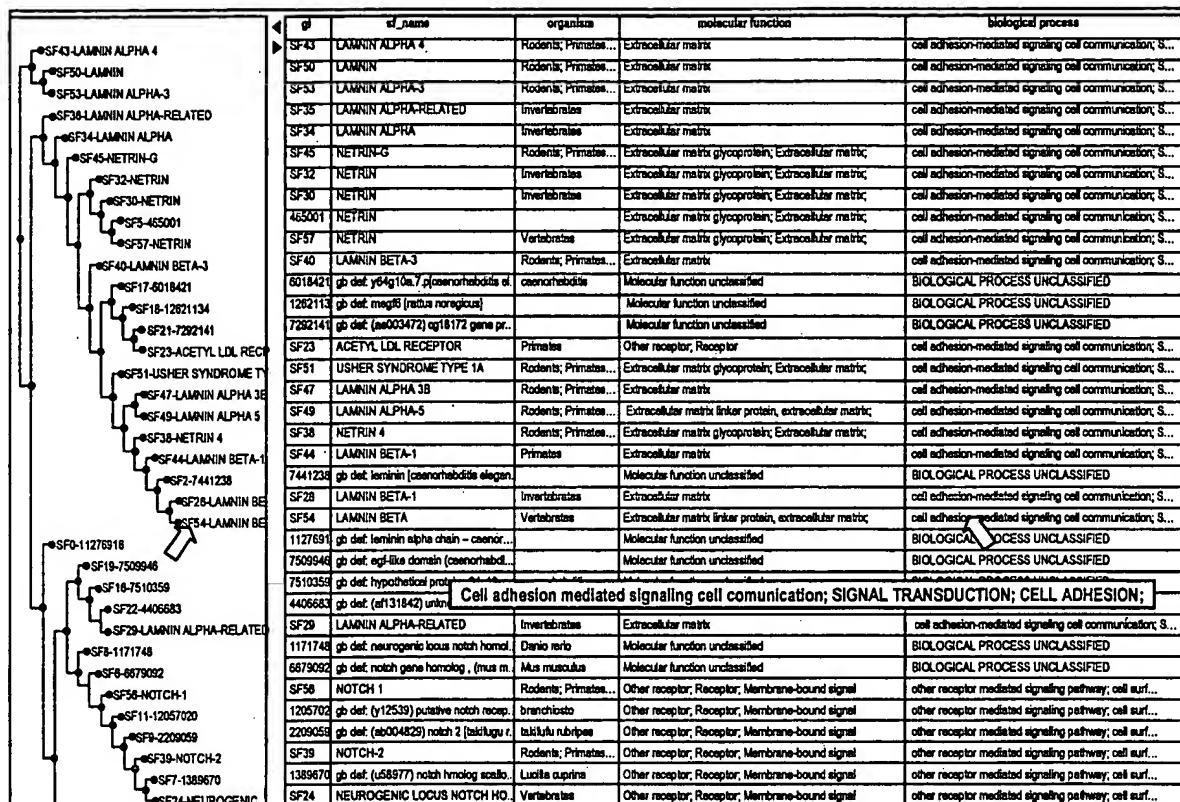
gi	sf_name	organism	molecular function	biological process
SF0	APOLIPOPROTEIN E	Mammals,Rodents,Primates...	Apolipoprotein,transfer carrier protein	lipid and fatty acid transport, LIPID, FATTY ACID AND STEROID METABOLISM, lipid and fa
SF1	APOLIPOPROTEIN A-IV	Mammals,Rodents,Primates...	Apolipoprotein,transfer carrier protein	lipid and fatty acid transport, LIPID, FATTY ACID AND STEROID METABOLISM, lipid and fa
SF2	APOLIPOPROTEIN A-1	Mammals,Rodents,Primates...	Apolipoprotein,transfer carrier protein	lipid and fatty acid transport, LIPID, FATTY ACID AND STEROID METABOLISM, lipid and fa
Lipid and fatty acid transport, LIPID, FATTY ACID AND STEROID METABOLISM, lipid and fatty acid transport, TRANSPORT, BLOOD CIRCULATION AND GAS EXCHANGE;				

**Fig. 21A**

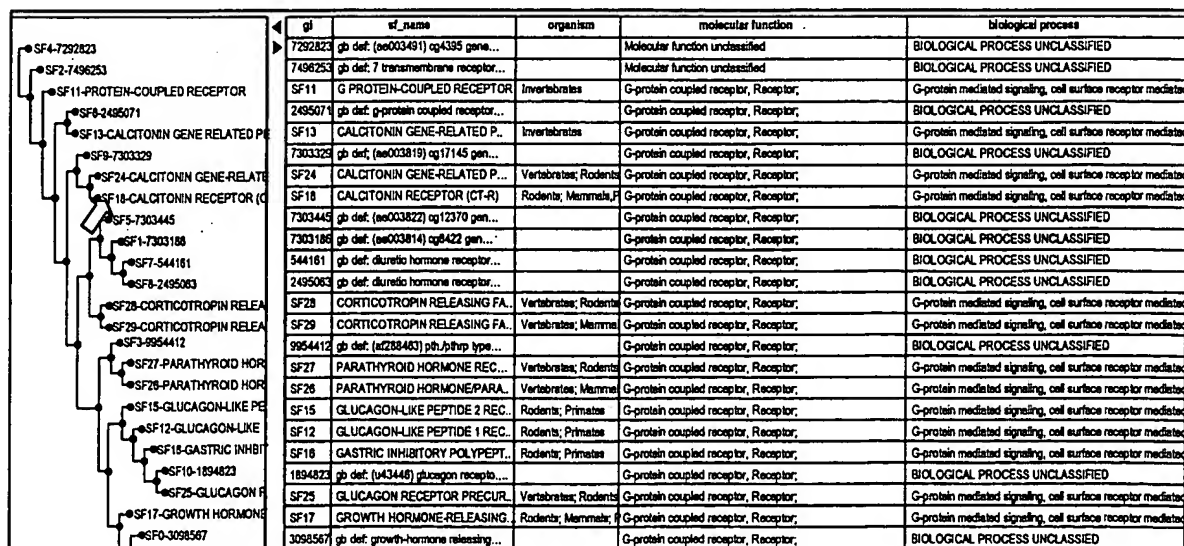
gi	sf_name	definition	organism
416828	APOLIPOPROTEIN E	APOLIPOPROTEIN E PRECURSOR (APO-E)	bos taurus
10644783	APOLIPOPROTEIN E	apolipoprotein E [Tupaia glis]	tupaia glis
3913071	APOLIPOPROTEIN E	APOLIPOPROTEIN E (APO-E)	saimiri sciureus
114040	APOLIPOPROTEIN E	APOLIPOPROTEIN E PRECURSOR (APO-E)	macaca fascicularis
114042	APOLIPOPROTEIN E	APOLIPOPROTEIN E PRECURSOR (APO-E)	papio hamadryas anubis
3913070	APOLIPOPROTEIN E	APOLIPOPROTEIN E PRECURSOR (APO-E)	macaca mulatta
11066430	APOLIPOPROTEIN E	apolipoprotein E [Hylobates lar]	hylobates lar
178853	APOLIPOPROTEIN E	apolipoprotein E	homo sapiens
4557325	APOLIPOPROTEIN E	APOLIPOPROTEIN E PRECURSOR (APO-E)	homo sapiens
178849	APOLIPOPROTEIN E	apolipoprotein E	homo sapiens
11066425	APOLIPOPROTEIN E	apolipoprotein E [Pongo pygmaeus]	pongo pygmaeus
11034803	APOLIPOPROTEIN E	apolipoprotein E [Pan troglodytes]	pan troglodytes
11066420	APOLIPOPROTEIN E	apolipoprotein E [Gorilla gorilla]	gorilla gorilla
114008	APOLIPOPROTEIN A-IV	APOLIPOPROTEIN A-IV PRECURSOR (APO-...	rattus norvegicus
8392909	APOLIPOPROTEIN A-IV	apolipoprotein C-IV [Rattus norvegicus]	rattus norvegicus
6880702	APOLIPOPROTEIN A-IV	apolipoprotein A-IV [Mus musculus]	mus musculus
191889	APOLIPOPROTEIN A-IV	apolipoprotein A-IV	mus musculus castaneus
12836356	APOLIPOPROTEIN A-IV	putative [Mus musculus]	mus musculus
1703331	APOLIPOPROTEIN A-IV	APOLIPOPROTEIN A-IV PRECURSOR (APO-...	mus musculus
109575	APOLIPOPROTEIN A-IV	apolipoprotein A-IV precursor - mouse (str...	mus musculus
3845997	APOLIPOPROTEIN A-IV	apolipoprotein AIV [Gallus gallus]	gallus gallus
3913046	APOLIPOPROTEIN A-IV	APOLIPOPROTEIN A-IV PRECURSOR (APO-...	sus scrofa
2492913	APOLIPOPROTEIN A-IV	APOLIPOPROTEIN A-IV PRECURSOR (APO-...	papio hamadryas anubis
481521	APOLIPOPROTEIN A-IV	APOLIPOPROTEIN A-IV PRECURSOR (APO-...	macaca fascicularis
71797	APOLIPOPROTEIN A-IV	apolipoprotein A-IV precursor - human	homo sapiens
114006	APOLIPOPROTEIN A-IV	APOLIPOPROTEIN A-IV PRECURSOR (APO-...	homo sapiens
4502151	APOLIPOPROTEIN A-IV	apolipoprotein A-IV precursor - [Homo sapien...	homo sapiens
11440019	APOLIPOPROTEIN A-IV	apolipoprotein A-IV precursor - [Homo sapien...	homo sapiens
6886379	APOLIPOPROTEIN A-IV	APOLIPOPROTEIN A-1 PRECURSOR	

**Fig. 21B**

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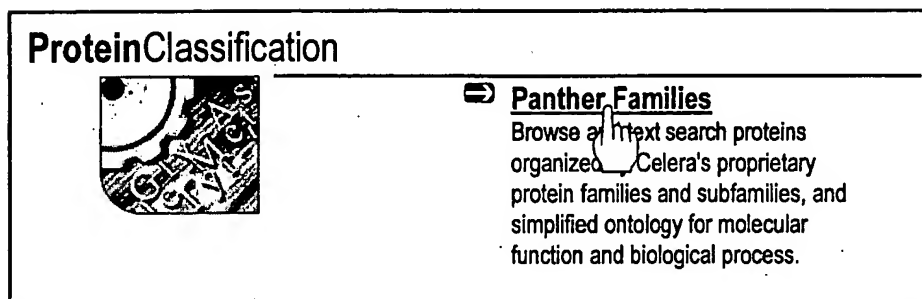


**Fig. 22A**







**Fig. 22B**

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**Fig. 23**

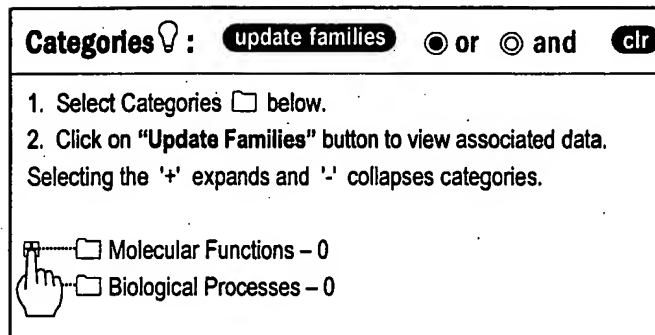
 <b>Panther Protein Function - Family Browser</b> <a href="#">Tips</a>		SEARCH  : <input type="radio"/> Categories <input type="radio"/> Families <input type="text"/> <input type="button" value="go"/>	
		SPECIES (gene list): <input checked="" type="checkbox"/> H. sapiens <input checked="" type="checkbox"/> M. musculus <input checked="" type="checkbox"/> D. melanogaster	
<b>Categories</b>  : <input type="button" value="update families"/> <input type="radio"/> or <input type="radio"/> and <input type="button" value="clr"/>		<b>Families</b>  : <input type="button" value="update categories"/> <input type="button" value="go to genelist"/> <input type="button" value="all"/> <input type="button" value="clr"/>	
<p>1. Select Categories <input type="checkbox"/> below.</p> <p>2. Click on "Update Families" button to view associated data.</p> <p>Selecting the '+' expands and '-' collapses categories.</p> <p><input type="checkbox"/> Molecular Functions - 0</p> <p><input type="checkbox"/> Biological Processes - 0</p>		<p>1. Select families <input type="checkbox"/> and or subfamilies <input type="checkbox"/> below.</p> <p>2. Click on "Go to Genelist" button to view associated genes.</p> <p>Highlighted subfamilies correspond to matches with your selected categories.</p>	

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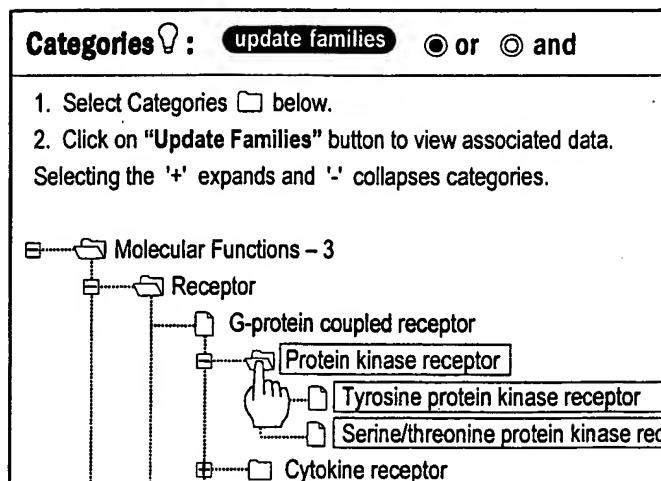
110

**Fig. 24**

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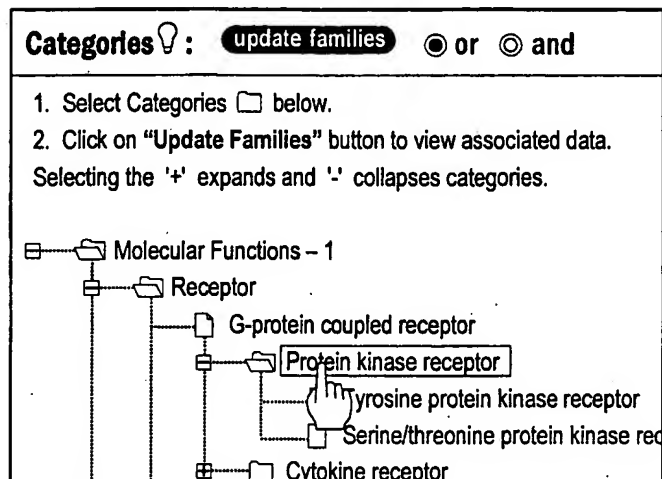


**Fig. 25**

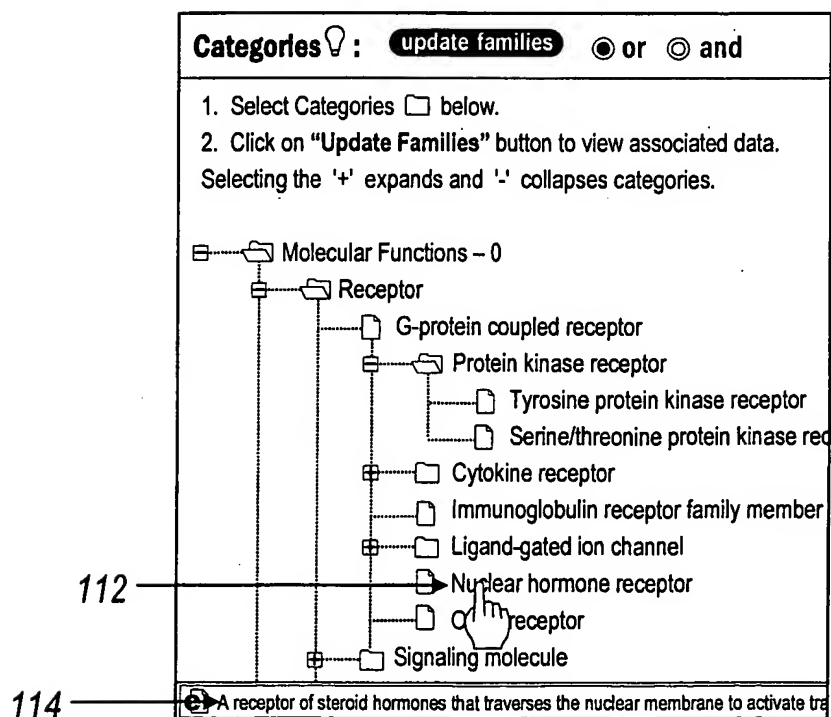


**Fig. 26**

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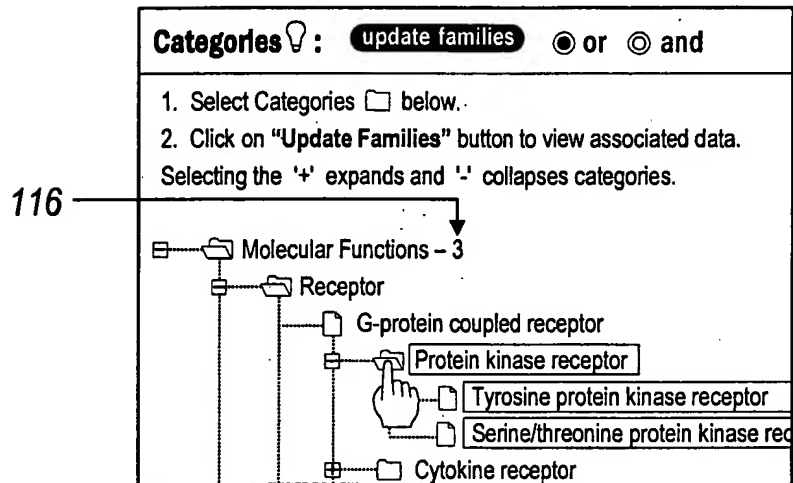


**Fig. 27**

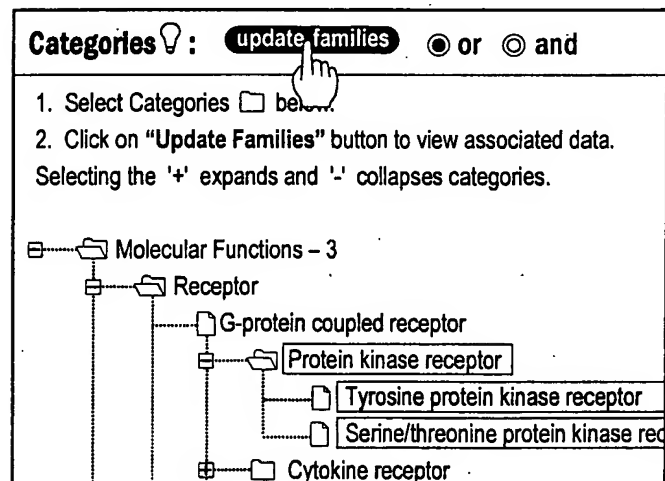


**Fig. 28**

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**Fig. 29**



**Fig. 30**

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Categories : <span style="background-color: black; color: white; padding: 2px 5px;">update families</span> <span style="border: 1px solid black; border-radius: 50%; padding: 0 5px;">or</span> <span style="border: 1px solid black; border-radius: 50%; padding: 0 5px;">and</span> <span style="background-color: black; color: white; padding: 2px 5px;">clr</span>	Families : <span style="background-color: black; color: white; padding: 2px 5px;">update categories</span> • <span style="background-color: black; color: white; padding: 2px 5px;">go to genelist</span>
<p>1. Select Categories <input type="checkbox"/> below.</p> <p>2. Click on "Update Families" button to view associated data.</p> <p>Selecting the '+' expands and '-' collapses categories.</p> <div style="margin-top: 10px;"> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;">Molecular Functions - 3</div> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;"> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;">Receptor</div> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;"> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;">G-protein coupled receptor</div> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;"> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;">Protein kinase receptor</div> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;">Tyrosine protein kinase recept</div> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;">Serine/threonine protein kinas</div> </div> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;">Cytokine receptor</div> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;">Immunoglobulin receptor family mem</div> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;">Ligand-gated ion channel</div> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;">Nuclear hormone receptor</div> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;">Other receptor</div> </div> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;">Signaling molecule</div> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;">Kinase</div> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;">Phosphatase</div> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;">Plateau</div> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;">Select regulatory molecule</div> </div> </div>	

**Fig. 31**

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Families : <span style="background-color: black; color: white; padding: 2px 5px;">update categories</span> • <span style="background-color: black; color: white; padding: 2px 5px;">go to genelist</span>
<p>1. Select families <input type="checkbox"/> and subfamilies <input type="checkbox"/> below.</p> <p>2. Click on "Go to Genelist" button to view associated genes.</p> <p>Highlighted subfamilies correspond to matches with your selected categories.</p>
<div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;"> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;">C-TYPE LECTIN-RELATED FAMILY MEMBER CF10148 - (1/20)</div> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;">Family tree   Full MSA   Partial MSA</div> </div> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;"> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;">SERINE/THREONINE PROTEIN KINASE CF 10467 - (3/10)</div> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;">Family tree   Full MSA   Partial MSA</div> </div> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;"> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;">TYROSINE-PROTEIN KINASE RECEPTOR CF 10301 - (13/13)</div> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;">Family tree   Full MSA   Partial MSA</div> </div> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;"> <div style="border: 1px solid black; padding: 2px; margin-bottom: 5px;">CELL ADHESION MOLECULE-RELATED CF11563 - (9/184)</div> </div>

**Fig. 32**

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<b>Families</b> : <a href="#">update categories</a> • <a href="#">go to genelist</a>		
1. Select families <input type="checkbox"/> and or subfamilies <input type="checkbox"/> below. 2. Click on "Go to Genelist" button to view associated genes. Highlighted subfamilies correspond to matches with your selected categories.		
<input checked="" type="checkbox"/>	<input type="checkbox"/>	C-TYPE LECTIN-RELATED FAMILY MEMBER CF10148 – (1/20)
Family tree	Full MSA	Partial MSA
<input checked="" type="checkbox"/>	<input type="checkbox"/>	SERINE/THREONINE PROTEIN KINASE CF 10467 – (3/10)
Family tree	Full MSA	Partial MSA
<input checked="" type="checkbox"/>	<input type="checkbox"/>	TYROSINE-PROTEIN KINASE RECEPTOR CF 10301 – (13/13)
Family tree	Full MSA	Partial MSA
<input checked="" type="checkbox"/>	<input type="checkbox"/>	CELL ADHESION MOLECULE-RELATED CF11563 – (9/184)

**Fig. 33**

<b>Families</b> : <a href="#">update categories</a> • <a href="#">go to genelist</a>		
1. Select families <input type="checkbox"/> and or subfamilies <input type="checkbox"/> below. 2. Click on "Go to Genelist" button to view associated genes. Highlighted subfamilies correspond to matches with your selected categories.		
<input checked="" type="checkbox"/>	<input type="checkbox"/>	C-TYPE LECTIN-RELATED FAMILY MEMBER CF10148 – (1/20)
Family tree	Full MSA	Partial MSA
<input checked="" type="checkbox"/>	<input type="checkbox"/>	SERINE/THREONINE PROTEIN KINASE CF 10467 – (3/10)
Family tree	Full MSA	Partial MSA
<input checked="" type="checkbox"/>	<input type="checkbox"/>	TYROSINE-PROTEIN KINASE RECEPTOR CF 10301 – (13/13)
Family tree	Full MSA	Partial MSA
<input checked="" type="checkbox"/>	<input type="checkbox"/>	CELL ADHESION MOLECULE-RELATED CF11563 – (9/184)

**Fig. 34**



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<b>Families</b> : <a href="#">update categories</a> • <a href="#">go to genelist</a>		
1. Select families <input type="checkbox"/> and or subfamilies <input type="checkbox"/> below. 2. Click on "Go to Genelist" button to view associated genes. Highlighted subfamilies correspond to matches with your selected categories.		
<input checked="" type="checkbox"/>	<input type="checkbox"/>	C-TYPE LECTIN-RELATED FAMILY MEMBER CF10148 – (1/20)
Family tree	Full MSA	Partial MSA
<input checked="" type="checkbox"/>	<input type="checkbox"/>	SERINE/THREONINE PROTEIN KINASE CF 10467 – (3/10)
Family tree	Full MSA	Partial MSA
<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	TYROSINE-PROTEIN KINASE RECEPTOR CF 10301 – (13/13)
Family tree	Full MSA	Partial MSA
<input checked="" type="checkbox"/>	<input type="checkbox"/>	CELL ADHESION MOLECULE-RELATED CF11563 – (9/184)

**Fig. 35**

<b>Families</b> : <a href="#">update categories</a> • <a href="#">go to genelist</a>		
1. Select families <input type="checkbox"/> and or subfamilies <input type="checkbox"/> below. 2. Click on "Go to Genelist" button to view associated genes. Highlighted subfamilies correspond to matches with your selected categories.		
<input checked="" type="checkbox"/>	<input type="checkbox"/>	C-TYPE LECTIN-RELATED FAMILY MEMBER CF10148 – (1/20)
Family tree	Full MSA	Partial MSA
<input checked="" type="checkbox"/>	<input type="checkbox"/>	SERINE/THREONINE PROTEIN KINASE CF 10467 – (3/10)
Family tree	Full MSA	Partial MSA
<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	TYROSINE-PROTEIN KINASE RECEPTOR CF 10301 – (13/13)
Family tree	Full MSA	Partial MSA
<input checked="" type="checkbox"/>	<input type="checkbox"/>	CELL ADHESION MOLECULE-RELATED CF11563 – (9/184)


**Fig. 36**

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### Protein Classification

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**Panther Families**  
 Browse and next search proteins organized by Celera's proprietary protein families and subfamilies.


**Fig. 37**

 <b>Panther Protein Function - Family Browser</b> <a href="#">Tips</a>		SEARCH  : <input type="radio"/> Categories <input type="radio"/> Families <input type="text"/> <input type="button" value="go"/>	
		SPECIES (gene list): <input checked="" type="checkbox"/> H. sapiens <input checked="" type="checkbox"/> M. musculus <input checked="" type="checkbox"/> D. melanogaster	
<b>Categories</b>  : <input type="button" value="update families"/> <input type="radio"/> or <input type="radio"/> and <input type="button" value="clr"/>		<b>Families</b>  : <input type="button" value="update categories"/> • <input type="button" value="go to genelist"/> <input type="button" value="all"/> <input type="button" value="clr"/>	
1. Select Categories <input type="checkbox"/> below. 2. Click on "Update Families" button to view associated data. Selecting the '+' expands and '-' collapses categories.  <div> <input type="checkbox"/> Molecular Functions - 0  <input type="checkbox"/> Biological Processes - 0                 </div>		1. Select families <input type="checkbox"/> and or subfamilies <input type="checkbox"/> below. 2. Click on "Go to Genelist" button to view associated genes. Highlighted subfamilies correspond to matches with your selected categories.	


**Fig. 38**


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SEARCH  : ☒ Categories ☐ Families



**Fig. 39**

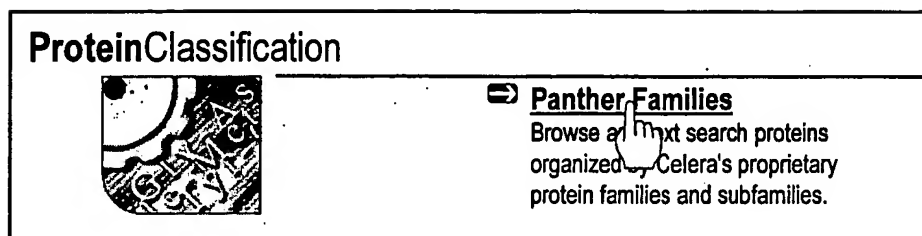
Categories  :  ☒ or ☐ and

1. Select Categories  below.  
2. Click on "Update Families" button to view associated data.  
Selecting the '+' expands and '-' collapses categories.


-  Protein kinase receptor
  -  Tyrosine protein kinase receptor
  -  Serine/threonine protein kinase receptor
-  Cytokine receptor
-  Immunoglobulin receptor family member
-  Ligand-gated ion channel
-  Nuclear hormone receptor
-  Other receptor
-  Signaling molecule
  -  Kinase
    -  Protein kinase
      -  Tyrosine protein kinase receptor
      -  Non-receptor tyrosine protein kinase
      -  Serine/threonine protein kinase receptor
      -  Non-receptor serine/threonine protein kinase
    -  Carbohydrate kinase
    -  Nucleotide kinase
    -  Amino acid kinase
    -  Other kinase

**Fig. 40**

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**Fig. 41**

 <b>Panther Protein Function - Family Browser</b> <a href="#">Tips</a>		SEARCH ? : <input type="radio"/> Categories <input type="radio"/> Families <input type="text"/> <input type="button" value="go"/>	
		SPECIES (gene list): <input checked="" type="checkbox"/> H. sapiens <input checked="" type="checkbox"/> M. musculus <input checked="" type="checkbox"/> D. melanogaster	
<b>Categories ? :</b> <input type="button" value="update families"/> <input type="radio"/> or <input type="radio"/> and <input type="button" value="clr"/>	<b>Families ? :</b> <input type="button" value="update categories"/> <input type="button" value="go to genelist"/> <input type="button" value="all"/> <input type="button" value="clr"/>		
1. Select Categories <input type="checkbox"/> below. 2. Click on "Update Families" button to view associated data. Selecting the '+' expands and '-' collapses categories.  <input type="checkbox"/> Molecular Functions - 0 <input type="checkbox"/> Biological Processes - 0		1. Select families <input type="checkbox"/> and or subfamilies <input type="checkbox"/> below. 2. Click on "Go to Genelist" button to view associated genes. Highlighted subfamilies correspond to matches with your selected categories.	


**Fig. 42**

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SEARCH  : ☐ Categories ☒ Families



**Fig. 43**

Families  :  •

1. Select Families ☐ and or subfamilies ☐ below.  
2. Click on "Go to Genelist" button to view associated data.  
Highlighted subfamilies correspond to searches with your selected categories

<input checked="" type="checkbox"/>	T-CELL RECEPTOR BETA CHAIN CF10022 (3/7)	<input type="checkbox"/>
Family tree	Full MSA Partial MSA	
<input checked="" type="checkbox"/>	T-CELL RECEPTOR BETA CHAIN CF10433 (2/2)	<input type="checkbox"/>
Family tree	Full MSA Partial MSA	
<input checked="" type="checkbox"/>	T-CELL RECEPTOR BETA CHAIN CF10450 (3/3)	<input type="checkbox"/>
Family tree	Full MSA Partial MSA	
<input checked="" type="checkbox"/>	T-CELL RECEPTOR BETA CHAIN CF10452 (1/1)	<input type="checkbox"/>
Family tree	Full MSA Partial MSA	
<input checked="" type="checkbox"/>	T-CELL RECEPTOR ALPHA CHAIN CF10475 (10/10)	<input type="checkbox"/>
Family tree	Full MSA Partial MSA	
<input checked="" type="checkbox"/>	T-CELL RECEPTOR BETA CHAIN CF10481 (5/7)	<input type="checkbox"/>
Family tree	Full MSA Partial MSA	
<input checked="" type="checkbox"/>	T-CELL RECEPTOR BETA CHAIN CF10610 (6/7)	<input type="checkbox"/>
Family tree	Full MSA Partial MSA	
<input checked="" type="checkbox"/>	T-CELL RECEPTOR DELTA CHAIN CF10850 (3/3)	<input type="checkbox"/>
Family tree	Full MSA Partial MSA	
<input checked="" type="checkbox"/>	T-CELL RECEPTOR BETA CHAIN CF10930 (1/1)	<input type="checkbox"/>
Family tree	Full MSA Partial MSA	
<input checked="" type="checkbox"/>	T-CELL RECEPTOR BETA CHAIN CF11424 (7/12)	<input type="checkbox"/>
Family tree	Full MSA Partial MSA	
<input checked="" type="checkbox"/>	T-CELL RECEPTOR GAMMA CHAIN CF11488 (8/7)	<input type="checkbox"/>
Family tree	Full MSA Partial MSA	
<input checked="" type="checkbox"/>	T-CELL RECEPTOR BETA CHAIN CF10477 (2/2)	<input type="checkbox"/>
Family tree	Full MSA Partial MSA	
<input checked="" type="checkbox"/>	T-CELL RECEPTOR BETA CHAIN CF10449 (4/4)	<input type="checkbox"/>
Family tree	Full MSA Partial MSA	
<input checked="" type="checkbox"/>	T-CELL RECEPTOR BETA CHAIN CF10451 (2/2)	<input type="checkbox"/>
Family tree	Full MSA Partial MSA	

**Fig. 44**


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Families


1. Select Families ☐ and or subfamilies ☐ below.  
2. Click on "Go to Genelist" button to view associated data.  
Highlighted subfamilies correspond to searches with your selected categories

☐ C-TYPE LECTIN-RELATED FAMILY MEMBER CF10148 (1/20)  
Family tree Full MSA Partial MSA

**Fig. 45**



Function Family Browser: Protein Families



HMM Score Cutoff (<):

-35.0

Display:

20

update

export

Hits 1-10 of 5414

[ page: (1) 2 3 4 5 6 ]

Sort results by selecting column title. Columns will sort in descending or ascending order.

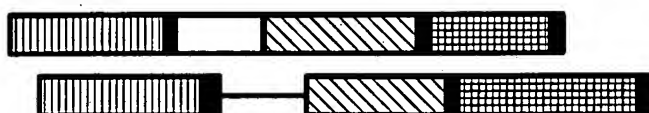
<div> <div>clr</div> <div>all</div> </div> <div>ID-protein</div> <div>Panther Best Hit – Panther ID family (CF#) or subfamily (SF#)</div> <div>Panther Score/Relation</div> <div>Species</div>					
<input type="checkbox"/>	1. <a href="#">hCP43060</a>	<a href="#">BONE MORPHOGENETIC PROTEIN RECEPTOR TYPE 1A (CF1...</a>	<a href="#">-1071.90</a>	***	<i>H. sapiens</i>
<input type="checkbox"/>	2. <a href="#">hCP20073</a>	<a href="#">ACTIVIN RECEPTOR TYPE I (CF11901-SF30)</a>	<a href="#">-1068.05</a>	***	<i>M. musculus</i>
<input type="checkbox"/>	3. <a href="#">hCP5773</a>	<a href="#">TGF-BETA RECEPTOR TYPE I (CF11901-SF28)</a>	<a href="#">-1046.96</a>	***	<i>M. musculus</i>
<input type="checkbox"/>	4. <a href="#">hCP44799</a>	<a href="#">TGF-BETA RECEPTOR TYPE I (CF11901-SF26)</a>	<a href="#">-1043.75</a>	***	<i>H. sapiens</i>
<input type="checkbox"/>	5. <a href="#">hCP9004</a>	<a href="#">SERINE/THIONINE-PROTEIN KINASE RECEPTOR R3 (CF1190...</a>	<a href="#">-1004.39</a>	***	<i>M. musculus</i>
<input type="checkbox"/>	6. <a href="#">hCP18779</a>	<a href="#">TYROSINE-PROTEIN KINASE 7-RELATED (CF11901-SF167)</a>	<a href="#">-1004.39</a>	***	<i>M. musculus</i>
<input type="checkbox"/>	7. <a href="#">hCP20049</a>	<a href="#">ACTIVIN RECEPTOR-LINE KINASE 7 (CF11901-SF20)</a>	<a href="#">-1024.83</a>	***	<i>M. musculus</i>
<input type="checkbox"/>	8. <a href="#">hCP18624</a>	<a href="#">TGF-BETA RECEPTOR TYPE II (CF11901-SF22)</a>	<a href="#">-1002.02</a>	***	<i>M. musculus</i>
<input type="checkbox"/>	9. <a href="#">hCP20668</a>	<a href="#">BONE MORPHOGENETIC PROTEIN RECEPTOR TYPE IA (CF1...</a>	<a href="#">-864.3</a>	***	<i>M. musculus</i>
<input type="checkbox"/>	10. <a href="#">hCP51898</a>	<a href="#">ACTIVIN RECEPTOR TYPE IB (CF11901-SF28)</a>	<a href="#">-980.95</a>	***	<i>H. sapiens</i>
<input type="checkbox"/>	11. <a href="#">hCP39834</a>	<a href="#">ACTIVIN RECEPTOR TYPE II (CF11901-SF21)</a>	<a href="#">-689.85</a>	***	<i>H. sapiens</i>

**Fig. 46**

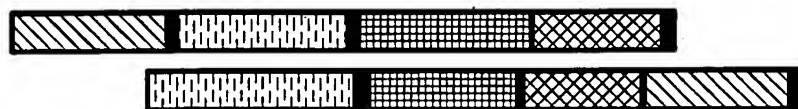
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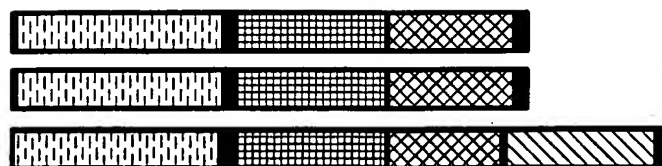
**FAMILY A**



**FAMILY B**



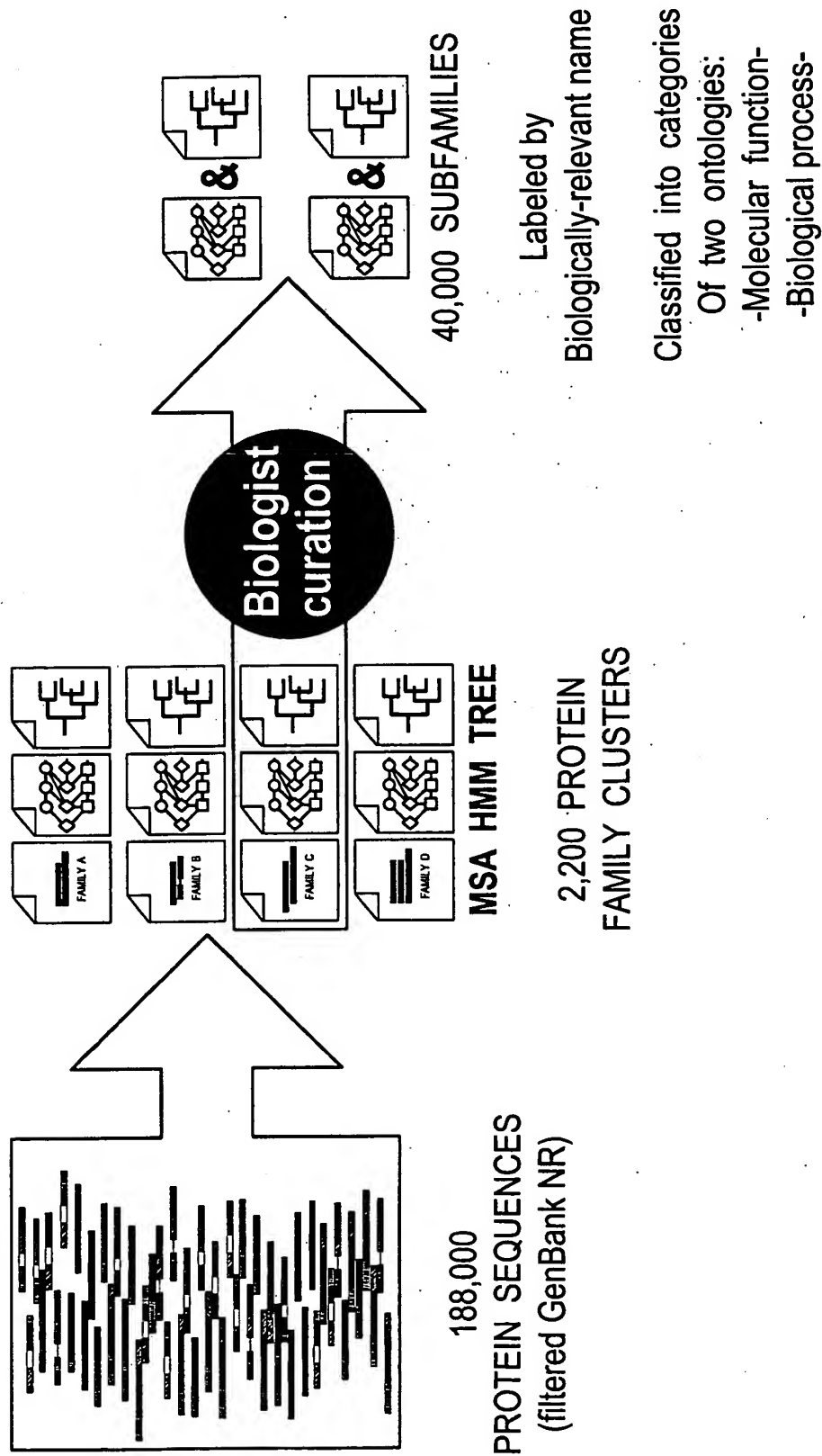
**FAMILY C**



**FAMILY D**

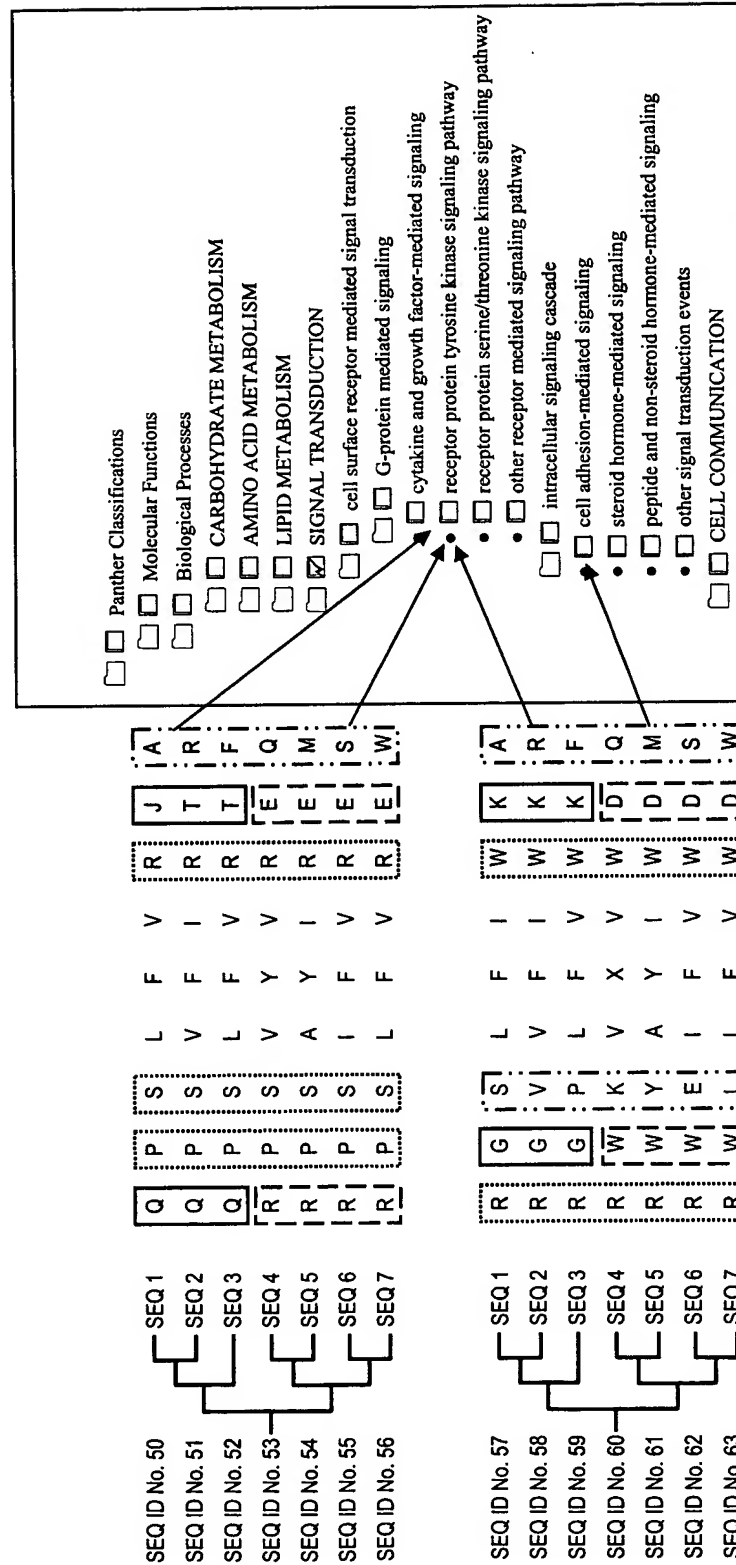
***Fig. 47***

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**Fig. 48**





**Fig. 49**